MLLNVLRICI	IVCLVNDGAG	KHSEGRERTK	TYSLNSRGYF	40
RKERGARRSK	ILLVNTKGLD	EPHIGHGDFG	LVAELFDSTR	80
THTNRKEPDM	NKVKLFSTVA	HGNKSARRKA	YNGSRRNIFS	120
RRSFDKRNTE	VTEKPGAKMF	WNNFLVKMNG	APQ <u>NTS</u> HGSK	160
AQEIMKEAC K	TLPFTQNIVH	ENCDRMVIQN	NLCFGKCISL	200
HVPNQQDRRN	TCSHCLPSKF	TLNHLTLNCT	GSKNVVKVVM	240
MVEECTCEAH	KSNFHQTAQF	NMDTSTTLHH		270

Figure 1. Deduced amino acid sequence of Xenopus cerberus protein. SEQ ID NO:1.

Figure 2. Nucleotide sequence of the full-length cerberus DNA derived from the Xenopus organizer. The sense strand is on top (in the 5' to 3' direction) and the antisense strand on the bottom line (on the opposite direction). SEQ ID NO:2.

GAATTCCCAG	CAAGTCGCTC	AGAAACACTG	CAGGGTCTAG	ATATCATACA	ATGTTACTAA	60
CTTAAGGGTC	GTTCAGCGAG	TCTTTGTGAC	GTCCCAGATC	TATAGTATGT	TACAATGATT	
ATGTACTCAG	GATCTGTATT	ATCGTCTGCC	TTGTGAATGA	TGGAGCAGGA	AAACACTCAG	120
TACATGAGTC	CTAGACATAA	TAGCAGACGG	AACACTTACT	ACCTCGTCCT	TTTGTGAGTC	
AAGGACGAGA	AAGGACAAAA	ACATATTCAC	TTAACAGCAG	AGGTTACTTC	AGAAAAGAAA	180
TTCCTGCTCT	TTCCTGTTTT	TGTATAAGTG	AATTGTCGTC	TCCAATGAAG	TCTTTTCTTT	
6166166166						
GAGGAGCACG	TAGGAGCAAG	ATTCTGCTGG	TGAATACTAA	AGGTCTTGAT	GAACCCCACA	240
CTCCTCGTGC	ATCCTCGTTC	TAAGACGACC	ACTTATGATT	TCCAGAACTA	CTTGGGGTGT	
MMCCCCO MCC	MC3.MMMM.000					
AACCCCTACC	TGATTTTCGC	TTAGTAGCTG	AACTATTTGA	TTCCACCAGA	ACACATACAA	300
AACCCGIACC	ACTAAAAGCG	AATCATCGAC	TTGATAAACT	AAGGTGGTCT	TGTGTATGTT	
ACAGAAAAGA	CCCACACAMO	1101110000				
TCTCTTTTCT	GCCAGACATG	MACAAAGTCA	AGCTTTTCTC	AACAGTTGCC	CATGGAAACA	360
1910111101	CGGTCTGTAC	TIGITICAGT	TCGAAAAGAG	TTGTCAACGG	GTACCTTTGT	
AAAGTGCAAG	AAGAAAAGCT	ጥ እ	C#1 C11 CC11			
TTTCACGTTC	TTCTTTTCGA	ATCTTACCAA	CINGANGGAA	TATTTTTCCT	CGCCGTTCTT	420
		HIGI INCOM	GATCTTCCTT	ATAAAAAGGA	GCGGCAAGAA	
TTGATAAAAG	AAATACAGAG	GTTACTCAAA	ACCCTCCTCC	C3.3.C3.TCTTC	#CC33C33###	400
AACTATTTTC	TTTATGTCTC	CAATGACTTT	TOGGACCACG	GTTCTACAAC	ACCERCARII	480
				GIICINCANG	ACCITGITAN	
TTTTGGTTAA	AATGAATGGA	GCCCCACAGA	ATACAAGCCA	TGGCAGTAAA	GCACAGGAAA	540
AAAACCAATT	TTACTTACCT	CGGGGTGTCT	TATGTTCGGT	ACCGTCATTT	CGTGTCCTTT	340
TAATGAAAGA	AGCTTGCAAA	ACCTTGTTTT	TCACTCAGAA	TATTGTACAT	GAAAACTGTG	600
ATTACTTTCT	TCGAACGTTT	TGGAACAAAA	AGTGAGTCTT	ATAACATGTA	CTTTTGACAC	
ACAGGATGGT	GATACAGAAC	AATCTGTGCT	TTGGTAAATG	CATCTCTCTC	CATGTTCCAA	660
TGTCCTACCA	CTATGTCTTG	TTAGACACGA	AACCATTTAC	GTAGAGAGAG	GTACAAGGTT	
1001001101	500100111					
MACAGCAAGA	TCGACGAAAT	ACTTGTTCCC	ATTGCTTGCC	GTCCAAATTT	ACCCTGAACC	720
TAGICGITCT	AGCTGCTTTA	TGAACAAGGG	TAACGAACGG	CAGGTTTAAA	TGGGACTTGG	
ACCTCACCCT	C N N M M C M N C M	CC1=c=1.				
TGGACTGCCA	GAATTGTACT	GGATCTAAGA	ATGTAGTAAA	GGTTGTCATG	ATGGTAGAGG	780
100AC1GCGA	CTTAACATGA	CCTAGATTCT	TACATCATTT	CCAACAGTAC	TACCATCTCC	
AATGCACGTG	TGAAGCTCAT	A B C B C C B B C M	6 001001110	50010105		
TTACGTGCAC	ACTTCGAGTA	TTCTCCTTCA	ACCECCERENC ACCECCERENC	ACCOCOCOLA	AACATGGATA	840
		110100110A	AGGIGGIIIG	ACGIGICAAA	TTGTACCTAT	
CATCTACTAC	CCTGCACCAT	TAAAGGACTG	CCATACAGTA	TEGALATECE	CTTTTTCTTCC	000
GTAGATGATG	GGACGTGGTA	ATTTCCTGAC	GGTATGTCAT	ACCTTTACGG	GAAAACAACC	900
AATATTTGTT	ACATACTATG	CATCTAAAGC	ATTATGTTGC	CTTCTATTTC	ATATAACCAC	960
TTATAAACAA	TGTATGATAC	GTAGATTTCG	TAATACAACG	GAAGATAAAG	TATATTGGTG	200
	•					
ATGGAATAAG	GATTGTATGA	ATTATAATTA	ACAAATGGCA	TTTTGTGTAA	CATGCAAGAT	1020
TACCTTATTC	CTAACATACT	TAATATTAAT	TGTTTACCGT	AAAACACATT	GTACGTTCTA	

CTCTGTTCCA	TCAGTTGCAA	GATAAAAGGC	AATATTTGTT	TGACTTTTTT	TCTACAAAAT	T080
GAGACAAGGT	AGTCAACGTT	CTATTTTCCG	TTATAAACAA	ACTGAAAAAA	AGATGTTTTA	
GAATACCCAA	ATATATGATA	AGATAATGGG	GTCAAAACTG	TTAAGGGGTA	ATGTAATAAT	1140
	TATATACTAT					2210
AGGGACTAAG	TTTGCCCAGG	AGCAGTGACC	CATAACAACC	AATCAGCAGG	TATGATTTAC	1200
TCCCTGATTC	AAACGGGTCC	TCGTCACTGG	GTATTGTTGG	TTAGTCGTCC	ATACTAAATG	
#00#03 00#0	mmm					
	TTTAAAAGCA					1260
ACCAGTGGAC	AAATTTTCGT	TTGTAGAATA	ACCAACGATA	CCCAATGACG	AAGACCCGTT	
AATGTGTGCC	TCATAGGGGG	GTTAGTGTGT	TGTGTACTGA	ATAAATTGTA	TTTATTTCAT	1320
	AGTATCCCCC					
TGTTACAAAA	AAAAAAA					
ACAATGTTTT	TTTTTTTT					

Fig. 2. (Continuation page 2, SEQ ID NO:2).

MSF	TRKVDSL	LLLAIPGLAL	LLLPNAYCAS	CEPVRIPMCK	SMPWNMTKMP	nhlhhstqan	60
AII	AIEQFEG	LLTTECSQDL	LFFLCAMYAP	ICTIDFQHEP	IKPCKSVCER	ARAGCEPILI	120
KYF	UHTWPESL	ACEELPVYDR	GVCISPEAIV	TVEQGTDSMP	DFSMDSNNGN	CGSGREHCKC	180
KPN	KATQKTY	LKNNYNYVIR	akvkevkvkc	HDATAIVEVK	EILKSSLVNI	PKDTVTLYTN	240
SGC	CLCPQLVA	NEEYIIMGYE	DKERTRLLLV	EGSLAEKWRD	RLAKKVKRWD	QKLRRPRKSK	300
DPI	/APIPNKN	SNSRQARS					

Figure 3. Deduced amino acid sequence of Xenopus frazzled protein. SEQ ID NO:3.

Figure 4. Nucleotide sequence of the full-length frazzled cDNA derived from the Xenopus organizer. The sense strand of the DNA on top (5' to 3' direction) and the antisense strand on the bottom line (opposite direction). SEQ ID NO:4.

GAATTCCCTT	TCACACAGGA	CTCCTGGCAG	AGGTGAATGG	TTAGCCCTAT	GGATTTGGTT	60
CTTAAGGGAA	AGTGTGTCCT	GAGGACCGTC	TCCACTTACC	AATCGGGATA	CCTAAACCAA	
TGTTGATTTT	GACACATGAT	TGATTGCTTT	CAGATAGGAT	TGAAGGACTT	GGATTTTTAT	120
ACAACTAAAA	CTGTGTACTA	ACTAACGAAA	GTCTATCCTA	ACTTCCTGAA	CCTAAAAATA	
Cm > 1 mm cm cc	10mmmm 1 1 1 =					
CTAATTCTGC	ACTITIAAAT	TATCTGAGTA	ATTGTTCATT	TTGTATTGGA	TGGGACTAAA	180
GATTANGACG	TGAAAATTTA	ATAGACTCAT	TAACAAGTAA	AACATAACCT	ACCCTGATTT	
GATAAACTTA	ACTCCTTGCT	ምምምር እ ር ምምር ር	CC242222C02	63.3.CC#CCCC	BC1 CBBCB1	
CTATTTGAAT	TGAGGAACGA	AAACTGAACG	CCUTATACIA	AMMCCA CCCC	TGAGTTGTAG	240
			GGINIIIGNI	ATTCCACCCC	ACTCAACATC	
TTGCTTTTAC	ATGTGCCCAG	ATTTTCCCTG	TATTCCCTGT	ATTCCCTCTA	AAGTAAGCCT	300
AACGAAAATG	TACACGGGTC	TAAAAGGGAC	ATAAGGGACA	TAAGGGAGAT	TTCATTCCCA	300
ACACATACAG	GTTGGGCAGA	ATAACAATGT	CTCGAACAAG	GAAAGTGGAC	TCATTACTGC	360
TGTGTATGTC	CAACCCGTCT	TATTGTTACA	GAGCTTGTTC	CTTTCACCTG	AGTAATGACG	
TACTGGCCAT	ACCTGGACTG	GCGCTTCTCT	TATTACCCAA	TGCTTACTGT	GCTTCGTGTG	420
ATGACCGGTA	TGGACCTGAC	CGCGAAGAGA	ATAATGGGTT	ACGAATGACA	CGAAGCACAC	
N.C.C.O.M.C.M.C.C.C.	01m00001m0					
*CCCACACCC	GATCCCCATG	TGCAAATCTA	TGCCATGGAA	CATGACCAAG	ATGCCCAACC	480
1 COGACACOC	CTAGGGGTAC	ACGTTTAGAT	ACGGTACCTT	GTACTGGTTC	TACGGGTTGG	
ATCTCCACCA	CAGCACTCAA	GCC A A TGCCC A	かたいかたことも 3 m	mc3 3 C3 Cmmm	C1100mmno	
TAGAGGTGGT	GTCGTGAGTT	CGGTTACGGT	AGGACCGTTA	TGAACAGTTT ACTTCTCA AA	GAAGGTTTGC	540
		00011110001	AGGACCGIIA	ACTIGICAAA	CTTCCAAACG	
TGACCACTGA	ATGTAGCCAG	GACCTTTTGT	TCTTTCTGTG	TGCCATGTAT	CCCCCC 3 で でで	600
ACTGGTGACT	TACATCGGTC	CTGGAAAACA	AGAAAGACAC	ACGGTACATA	CGGGGGTAAA	800
GTACCATCGA	TTTCCAGCAT	GAACCAATTA	AGCCTTGCAA	GTCCGTGTGC	GAAAGGGCCA	660
CATGGTAGCT	AAAGGTCGTA	CTTGGTTAAT	TCGGAACGTT	CAGGCACACG	CTTTCCCGGT	
GGGCCGGCTG	TGAGCCCATT	CTCATAAAGT	ACCGGCACAC	TTGGCCAGAG	AGCCTGGCAT	720
CCCGGCCGAC	ACTCGGGTAA	GAGTATTTCA	TEGCCETETE	AACCGGTCTC	TCGGACCGTA	
GTG A A G A G C TT	CCCCCTATATA	0101010010				
CACTTCTCCA	GCCCGTATAT	GACAGAGGAG	TCTGCATCTC	CCCAGAGGCT	ATCGTCACAG	780
CHOI TO TOGA	CGGGCATATA	CIGICICCIC	AGACGTAGAG	GGGTCTCCGA	TAGCAGTGTC	
TGGAACAAGG	AACAGATTCA	ATGCCAGACT	ずごがごごみ かごごみ	*******	CC111mmccc	
ACCTTGTTCC	TTGTCTAAGT	TACGGTCTGA	AGAGGTACCT	DAGTTTGTTA	CCTTTARCC	840
				MOIIIGIIA	CCITIAACGC	
GAAGCGGCAG	GGAGCACTGT	AAATGCAAGC	CCATGAAGGC	AACCCAAAAG	ACGTATCTCA	900
CTTCGCCGTC	CCTCGTGACA	TTTACGTTCG	GGTACTTCCG	TTGGGTTTTC	TGCATAGAGT	300
AGAATAATTA	CAATTATGTA	ATCAGAGCAA	aagtgaaaga	GGTGAAAGTG	AAATGCCACG	960
TCTTATTAAT	GTTAATACAT	TAGTCTCGTT	TTCACTTTCT	CCACTTTCAC	TTTACGGTGC	
ACCCA ACACC						
TCCCTTCTCC	AATTGTGGAA	GTAAAGGAGA	TTCTCAAGTC	TTCCCTAGTG	AACATTCCTA	1020
-000110100	TTAACACCTT	CATTTCCTCT	AAGAGTTCAG	AAGGGATCAC	TTGTAAGGAT	

*						
AAGACACAGT	GACACTGTAC	ACCAACTCAG	GCTGCTTGTG	CCCCCAGCTT	GTTGCCAATG	1080
TTCTGTGTCA	CTGTGACATG	TGGTTGAGTC	CGACGAACAC	GGGGGTCGAA	CAACGGTTAC	
AGGAATACAT	AATTATGGGC	TATGAAGACA	AAGAGCGTAC	CAGGCTTCTA	CTAGTGGAAG	1140
TCCTTATGTA	TTAATACCCG	ATACTTCTGT	TTCTCGCATG	GTCCGAAGAT	GATCACCTTC	
		•				
GATCCTTGGC	CGAAAAATGG	AGAGATCGTC	TTGCTAAGAA	AGTCAAGCGC	TGGGATCAAA	1200
CTAGGAACCG	GCTTTTTACC	TCTCTAGCAG	AACGATTCTT	TCAGTTCGCG	ACCCTAGTTT	
•						
AGCTTCGACG	TCCCAGGAAA	AGCAAAGACC	CCGTGGCTCC	AATTCCCAAC	AAAAACAGCA	1260
TCGAAGCTGC	AGGGTCCTTT	TCGTTTCTGG	GGCACCGAGG	TTAAGGGTTG	TTTTTGTCGT	
ATTCCAGACA	AGCGCGTAGT	TAGACTAACG	GAAAGGTGTA	TGGAAACTCT	ATGGACTTTG	1320
TAAGGTCTGT	TCGCGCATCA	ATCTGATTGC	CTTTCCACAT	ACCTTTGAGA	TACCTGAAAC	
AAACTAAGAT	TTGCATTGTT	GGAAGAGCAA	AAAAGAAATT	GCACTACAGC	ACGTTATATT	1380
			TTTTCTTTAA			
CTATTGTTTA	CTACAAGAAG	CTGGTTTAGT	TGATTGTAGT	TCTCCTTTCC	TTCTTTTTTT	1440
GATAACAAAT	GATGTTCTTC	GACCAAATCA	ACTAACATCA	AGAGGAAAGG	AAGAAAAAA	
TTATAACTAT	ATTTGCACGT	GTTCCCAGGC	AATTGTTTTA	TTCAACTTCC	AGTGACAGAG	1500
			TTAACAAAAT			
CAGTGACTGA	ATGTCTCAGC	CTAAAGAAGC	TCAATTCATT	TCTGATCAAC	TAATGGTGAC	1560
			AGTTAAGTAA			
AAGTGTTTGA	TACTTGGGGA	AAGTGAACTA	ATTGCAATGG	TAAATCAGAG	AAAAGTTGAC	1620
			TAACGTTACC			
CAATGTTGCT	TTTCCTGTAG	ATGAACAAGT	GAGAGATCAC	ATTTAAATGA	TGATCACTTT	1680
GTTACAACGA	AAAGGACATC	TACTTGTTCA	CTCTCTAGTG	TAAATTTACT	ACTAGTGAAA	
CCATTTAATA	CTTTCAGCAG	TTTTAGTTAG	ATGACATGTA	GGATGCACCT	AAATCTAAAT	1740
GGTAAATTAT	GAAAGTCGTC	AAAATCAATC	TACTGTACAT	CCTACGTGGA	TTTAGATTTA	
ATTTTATCAT	AAATGAAGAG	CTGGTTTAGA	CTGTATGGTC	ACTGTTGGGA	AGGTAAATGC	1800
TAAAATAGTA	TTTACTTCTC	GACCAAATCT	GACATACCAG	TGACAACCCT	TCCATTTACG	
CTACTTTGTC	AATTCTGTTT	TAAAAATTGC	CTAAATAAAT	ATTAAGTCCT	AAATAAAAA	1860
GATGAAACAG	TTAAGACAAA	ATTTTTAACG	GATTTATTTA	TAATTCAGGA	TTTATTTTTT	
AAAAAAAAA	AAAAA					
TTTTTTTTT	TTTTT					

Fig. 4. (Continuation page 2, SEQ ID NO:4).

60 MLLLFRAIPM LLIGLMVLQT DCEIAQYYID EEEPPGTVIA VLSQHSIFNT TDIPATNFRL MKQFNNSLIG VRESDGQLSI MERIDREQIC RQSLHCNLAL DVVSFSKGHF KLLNVKVEVR 120 DINDHSPHFP SEIMHVEVSE SSSVGTRIPL EIAIDEDVGS NSIQNFQISN NSHFSIDVLT 180 RADGVKYADL VLMRELDREI QPTYIMELLA MDGGVPSLSG TAVVNIRVLD FNDNSPVFER 240 STIAVDLVED APLGYLLLEL HATDDDEGVN GEIVYGFSTL ASQEVRQLFK INSRTGSVTL 300 EGGVDFETKO TYEFEVOAOD LGPNPLTATC KVTVHILDVN DNTPAITITP LTTVNAGVAY 360 IPETATKENF IALISTIDRA SGSNGOVRCT LYGHEHFKLQ QAYEDSYMIV TISTLDRENI 420 AAYSLTVVAE DIGFPSLKTK KYYTVKVSDE NDNAPVFSKP QYEASILENN APGSYITTVI 480 ARDSDSDONG KVNYRLVDAK VMGOSLTTFV SLDADSGVLR AVRSLDYEKL KQLDFEIEAA 540 DNGIPQLSTR VQLNLRIVDQ NDNCPVITNP LLNNGSGEVL LPISAPQNYL VFQLKAEDSD 600 EGHNSQLFYT ILRDPSRLFA INKESGEVFL KKQLNSDHSE DLSIVVAVYD LGRPSLSTNA 660 TVKFILTDSF PSNVEVVILQ PSAEEQHQID MSIIFIAVLA GGCALLLLAI FFVACTCKKK 720 AGEFKQVPEQ HGTCNEERLL STPSPQSVSS SLSQSESCQL SINTESENCS VSSNQEQHQQ 780 TGIKHSISVP SYHTSGWHLD NCAMSISGHS HMGHISTKVQ WAKEIVTSMT VTLILVENQK 840 RRALSSQCRH KPVLNTQMNQ QGSDMPITIS ATESTRVQKM GTAHCNMKRA IDCLTL

Figure 5. Deduced amino acid sequence of the Xenopus PAPC (paraxial protocadherin) protein. It encodes a member of the cadherin family of transmembrane proteins that has dorsalizing activity when constructs are injected into Xenopus embryos. SEQ ID NO:5.

Figure 6. Nucleotide sequence of the full-length PAPC cDNA derived from the Xenopus organizer. The sense strand of the DNA is shown in the top line (in the 5' to 3' direction), and the bottom line shows the antisense strand (opposite orientation). SEQ ID NO:6.

AGATGAACTC TCTACTTGAG		 		60
ACTGTTTCTA TGACAAAGAT		 		120
 CTTCAAGATG GAAGTTCTAC		 		180
 ACAAACAGAC TGTTTGTCTG	-	 		240
 AATTGCAGTG TTAACGTCAC		 		300
 CCGTCTAATG GGCAGATTAC				360
GAGCATCATG CTCGTAGTAC				420
 GGCTTTGGAT CCGAAACCTA		 		480
 GGTGAGAGAC CCACTCTCTG		 		540
 GTCTGAAAGT CAGACTTTCA		 		600
TGGGTCCAAC ACCCAGGTTG		 		6 60
GCTAACCAGA CGATTGGTCT		 		720
GGAAATCCAG CCTTTAGGTC				780
 ATCTGGTACT TAGACCATGA		 		840
TGAGAGAAGC ACTCTCTTCG				900
GGAGTTACAT CCTCAATGTA		 		960
CACTTTGGCA CTGAAACCGT			AACTCCAGAA TTGAGGTCTT	1020

		· · · · · · · · · · · · · · · · · · ·	CAAGCAGACT GTTCGTCTGA		1080
			TACTTGTAAA ATGAACATTT		1140
			TACCCCTCTG ATGGGGAGAC		1200
			GAACTTTATA CTTGAAATAT	• •	1260
			CTGTACTCTT GACATGAGAA		1320
			GATAGTTACC CTATCAATGG	•	1380
 			TGCAGAAGAC ACGTCTTCTG		1440
 			TGATGAGAAT ACTACTCTTA		1500
 •			AAATAATGCT TTTATTACGA		1560
 			AAATGGCAAA TTTACCGTTT		1620
			ATTTGTTTCT TAAACAAAGA		1680
			AAAACTTAAA TTTTGAATTT		1740
			CACTCGCGTT GTGAGCGCAA		1800
			TAATCCTCTT ATTAGGAGAA		1860
			CTATTTAGTT GATAAATCAA	TTCCAGCTCA AAGGTCGAGT	1920
			CTATACCATA GATATGGTAT	CTGAGAGATC GACTCTCTAG	1980
 			GTTCCTGAAA CAAGGACTTT	AAACAATTAA TTTGTTAATT	2040
			-	GGAAGACCTT CCTTCTGGAA	2100
	_		CTCTTTTCCT GAGAAAAGGA	TCTAACGTTG AGATTGCAAC	2160

Fig. 6. (Continuation page 2, SEQ ID NO:6).

	TTTGCAACCA AAACGTTGGT			2220
	GCTGGCTGGT CGACCGACCA			2280
-	AAAGAAAGCT TTTCTTTCGA	 		2340
	CCTGTTAAGC GGACAATTCG	 		2400
	CCAACTCTCC GGTTGAGAGG	 		2460
	TCAGCAAACA AGTCGTTTGT	 		2520
	CCTGGACAAT GGACCTGTTA			2580
	GGTACAGTGG CCATGTCACC			2640
	TCAGAAAAGA AGTCTTTTCT	 		2700
	GAATCAGCAG CTTAGTCGTC			2760
	GAAAATGGGA CTTTTACCCT	 		2820
	CCTGTATATT GGACATATAA			2880
	CTTAGAGACC GAATCTCTGG			2940
	GAAAGAGATT CTTTCTCTAA	 	AGAGATCGTC TCTCTAGCAG	3000
		 	 ATCCTTCAGA TAGGAAGTCT	3060
		 	GCAAGTGCTT CGTTCACGAA	3120
			GGGGAGACAC CCCCTCTGTG	3180
			ATTTTTTGTT	3240
			CTAACTAGCA GATTGATCGT	3300

Fig. 6. (Continuation page 3, SEQ ID NO:6).

				TGAAACAGCA ACTTTGTCGT		3360
				TGCCCTCTGT		3420
CTGGATTTCA	CCGGAAAAAT	GAAAATCGTC	GAGGACCCAG	ACGGGAGACA	CAATTAGTCG	
CCCTGGTCAA	GTCCTGAGTA	GGATCATGGC	GTTTTTATAT	GCATCTCACC	TACTTTGGAC	3480
				CGTAGAGTGG		
GTGATTTACA	CATAATAGGA	AACGCTTGGT	TTCAGTGAAG	TCTGTGTTGT	ATATATTCTG	3540
CACTAAATGT	GTATTATCCT	TTGCGAACCA	AAGTCACTTC	AGACACAACA	TATATAAGAC	
TTATATACAC	GCATTTTGTG	TTTGTGTATA	TATTTCAAGT	CCATTCAGAT	ATGTGTATAT	3600
AATATATGTG	CGTAAAACAC	AAACACATAT	ATAAAGTTCA	GGTAAGTCTA	TACACATATA	
AGTGCAGACC	TTGTAAATTA	AATATTCTGA	TACTTTTTCC	TCAATAAATA	TTTAAAT	
TCACGTCTGG	AACATTTAAT	TTATAAGACT	ATGAAAAAGG	AGTTATTTAT	AAATTTA	

Fig. 6. (Continuation page 4, SEQ ID NO:6).

MVCCGPGRML LGWAGLLVLA ALCLLQVPGA QAAACEPVRI PLCKSLPWNM TKMPNHLHHS 60

TQANAILAME QFEGLLGTHC SPDLLFFLCA MYAPICTIDF QHEPIKPCKS VCERARQGCE 120

PILIKYRHSW PESLACDELP VYDRGVCISP EAIVTADGAD FPMDSSTGHC RGASSERCKC 180

KPVRATQKTY FRNNYNYVIR AKVKEVKMKC HDVTAVVEVK EILKASLVNI PRDTVNLYTT 240

SGCLCPPLTV NEEYVIMGYE DEERSRLLLV EGSIAEKWKD RLGKKVKRWD MKLRHLGLGK 300

TDASDSTQNQ KSGRNSNPRP ARS.

Figure 7. Deduced amino acid sequence of mouse FRZB-1 protein. SEQ ID NO:7.

Figure 8. Nucleotide sequence of the full-length mouse FRZB-1 cDNA. SEQ ID NO:8. AAGCCTGGGA CCATGGTCTG CTGCGGCCCG GGACGGATGC TGCTAGGATG GGCCGGGTTG 60 TTCGGACCCT GGTACCAGAC GACGCCGGGC CCTGCCTACG ACGATCCTAC CCGGCCCAAC CTAGTCCTGG CTGCTCTCTG CCTGCTCCAG GTGCCCGGAG CTCAGGCTGC AGCCTGTGAG GATCAGGACC GACGAGACC GGACGAGGTC CACGGGCCTC GAGTCCGACG TCGGACACTC CCTGTCCGCA TCCCGCTGTG CAAGTCCCTT CCCTGGAACA TGACCAAGAT GCCCAACCAC 180 GGACAGGCGT AGGGCGACAC GTTCAGGGAA GGGACCTTGT ACTGGTTCTA CGGGTTGGTG CTGCACCACA GCACCCAGGC TAACGCCATC CTGGCCATGG AACAGTTCGA AGGGCTGCTG 240 GACGTGGTGT CGTGGGTCCG ATTGCGGTAG GACCGGTACC TTGTCAAGCT TCCCGACGAC GGCACCCACT GCAGCCCGGA TCTTCTCTTC TTCCTCTGTG CAATGTACGC ACCCATTTGC 300 CCGTGGGTGA CGTCGGGCCT AGAAGAGAAG AAGGAGACAC GTTACATGCG TGGGTAAACG ACCATCGACT TCCAGCACGA GCCCATCAAG CCCTGCAAGT CTGTGTGTGA GCGCGCCCGA TGGTAGCTGA AGGTCGTGCT CGGGTAGTTC GGGACGTTCA GACACACACT CGCGCGGCCT CAGGGCTGCG AGCCCATTCT CATCAAGTAC CGCCACTCGT GGCCGGAAAG CTTGGCCTGC 420 GTCCCGACGC TCGGGTAAGA GTAGTTCATG GCGGTGAGCA CCGGCCTTTC GAACCGGACG GACGAGCTGC CGGTGTACGA CCGCGGCGTG TGCATCTCTC CTGAGGCCAT CGTCACCGCG 480 CTGCTCGACG GCCACATGCT GGCGCCGCAC ACGTAGAGAG GACTCCGGTA GCAGTGGCGC GACGGAGCGG ATTTTCCTAT GGATTCAAGT ACTGGACACT GCAGAGGGGC AAGCAGCGAA 540 CTGCCTCGCC TAAAAGGATA CCTAAGTTCA TGACCTGTGA CGTCTCCCCG TTCGTCGCTT CGTTGCAAAT GTAAGCCTGT CAGAGCTACA CAGAAGACCT ATTTCCGGAA CAATTACAAC 600 GCAACGTTTA CATTCGGACA GTCTCGATGT GTCTTCTGGA TAAAGGCCTT GTTAATGTTG TATGTCATCC GGGCTAAAGT TAAAGAGGTA AAGATGAAAT GTCATGATGT GACCGCCGTT 660 ATACAGTAGG CCCGATTTCA ATTTCTCCAT TTCTACTTTA CAGTACTACA CTGGCGGCAA GTGGAAGTGA AGGAAATTCT AAAGGCATCA CTGGTAAACA TTCCAAGGGA CACCGTCAAT 720 CACCTTCACT TCCTTTAAGA TTTCCGTAGT GACCATTTGT AAGGTTCCCT GTGGCAGTTA CTTTATACCA CCTCTGGCTG CCTCTGTCCT CCACTTACTG TCAATGAGGA ATATGTCATC 780 GAAATATGGT GGAGACCGAC GGAGACAGGA GGTGAATGAC AGTTACTCCT TATACAGTAG ATGGGCTATG AAGACGAGGA ACGTTCCAGG TTACTCTTGG TAGAAGGCTC TATAGCTGAG 840 TACCCGATAC TTCTGCTCCT TGCAAGGTCC AATGAGAACC ATCTTCCGAG ATATCGACTC AAGTGGAAGG ATCGGCTTGG TAAGAAAGTC AAGCGCTGGG ATATGAAACT CCGACACCTT 900 TTCACCTTCC TAGCCGAACC ATTCTTTCAG TTCGCGACCC TATACTTTGA GGCTGTGGAA

GGACTGGGTA AAACTGATGC TAGCGATTCC ACTCAGAATC AGAAGTCTGG CAGGAACTCT

CCTGACCCAT TTTGACTACG ATCGCTAAGG TGAGTCTTAG TCTTCAGACC GTCCTTGAGA

960

AATCCCCGGC CAGCACGCAG CTAAATCCTG AAATGTAAAA GGCCACACCC ACGGAC TTAGGGGCCG GTCGTGCGTC GATTTAGGAC TTTACATTTT CCGGTGTGGG TGCCTG	
TTCTAAGACT GGCGCTGGTG GACTAACAAA GGAAAACCGC ACAGTTGTGC TCGTGAAAAAAAAAA	
TTGTTTACCG CAGACACCGC GTGGCTACCG AAGTTACTTC CGGTCCCCTT TCTCCTAACAAATGGC GTCTGTGGCG CACCGATGGC TTCAATGAAG GCCAGGGGAA AGAGGA	GCTT 1140 CGAA
CTTAATGGCG TGGGGTTAGA TCCTTTAATA TGTTATATAT TCTGTTTCAT CAATCA GAATTACCGC ACCCCAATCT AGGAAATTAT ACAATATATA AGACAAAGTA GTTAGT	CGTG 1200 CGCAC
GGGACTGTTC TTTTGCAACC AGAATAGTAA ATTAAATATG TTGATGCTAA GGTTTCCCTGACAAG AAAACGTTGG TCTTATCATT TAATTTATAC AACTACGATT CCAAAC	ETGTA 1260 BACAT
CTGGACTCCC TGGGTTTAAT TTGGTGTTCT GTACCCTGAT TGAGAATGCA ATGTTTGACCTGAGGG ACCCAAATTA AACCACAAGA CATGGGACTA ACTCTTACGT TACAAA	CATG 1320 AGTAC
TAAAGAGAGA ATCCTGGTCA TATCTCAAGA ACTAGATATT GCTGTAAGAC AGCCTCATTCTCTCT TAGGACCAGT ATAGAGTTCT TGATCTATAA CGACATTCTG TCGGAC	CTGCT 1380 GACGA
GCTGCGCTTA TAGTCTTGTG TTTGTATGCC TTTGTCCATT TCCCTCATGC TGTGA CGACGCGAAT ATCAGAACAC AAACATACGG AAACAGGTAA AGGGAGTACG ACACT	AAGTT 1440 ITCAA
ATACATGTTT ATAAAGGTAG AACGGCATTT TGAAATCAGA CACTGCACAA GCAGA TATGTACAAA TATTTCCATC TTGCCGTAAA ACTTTAGTCT GTGACGTGTT CGTCT	GTAGC 1500 CATCG
CCAACACCAG GAAGCATTTA TGAGGAAACG CCACACAGCA TGACTTATTT TCAAG GGTTGTGGTC CTTCGTAAAT ACTCCTTTGC GGTGTGTCGT ACTGAATAAA AGTTC	ATTGG 1560 TAACC
CAGGCAGCAA AATAAATAGT GTTGGGAGCC AAGAAAAGAA	AGGGG 1620 TCCCC
CACACTGGAA TCAGTAGCCC TTGAGCCATT AACAGCAGTG TTCTTCTGGC AAGTT GTGTGACCTT AGTCATCGGG AACTCGGTAA TTGTCGTCAC AAGAAGACCG TTCAA	TTTGA 1680 AAACT
TTTGTTCATA AATGTATTCA CGAGCATTAG AGATGAACTT ATAACTAGAC ATCTC AAACAAGTAT TTACATAAGT GCTCGTAATC TCTACTTGAA TATTGATCTG TAGAC	ETTGTT 1740 CAACAA
ATCTCTATAG CTCTGCTTCC TTCTAAATCA AACCCATTGT TGGATGCTCC CTCTC TAGAGATATC GAGACGAAGG AAGATTTAGT TTGGGTAACA ACCTACGAGG GAGAC	CCATTC 1800 GGTAAG

	TTGGCTTGCT AACCGAACGA			1860
	GTGTTATTTA CACAATAAAT			1920
	GTGCACATTT CACGTGTAAA			1980
	TGTGTTTATG ACACAAATAC			2040
	ACTAGATTAG TGATCTAATC			2100
	TAATGCTCCA ATTACGAGGT			2160
CGACAACAAC				

MVCGSPGGML	LLRAGLLALA	ALCLLRVPGA	RAAACEPVRI	PLCKSLPWNM	TKMPNHLHHS	60
TQANAILAIE	QFEGLLGTHC	SPDLLFFLCA	MYAPICTIDF	QHEPIKPCKS	VCERARQGCE	120
PILIKYRHSW	PENLACEELP	VYDRGVCISP	EAIVTADGAD	FPMDSSNGNC	RGASSERCKC	180
KPIRATQKTY	FRNNYNYVIR	AKVKEIKTKC	HDVTAVVEVK	EILKSSLVNI	PRDTVNLYTS	240
SGCLCPPLNV	NEEYIIMGYE	DEERSRLLLV	EGSIAEKWKD	RLGKKVKRWD	MKLRHLGLSK	300
SDSSNSDSTQ	SQKSGRNSNP	RQARN.				

Figure 9. Deduced amino acid sequence of human FRZB-1 protein. SEQ ID NO:9.

Figure 10. Nucleotide sequence of the full-length human FRZB-1 cDNA. SEQ ID NO:10. This sequence was assembled from public ESTs from the Genbank database (accession numbers: H18848, R63748, W38677, W44760, H38379 and N71244).

GGCGGAGCGG	GCCTTTTGGC	GTCCACTGCG (CGGCTGCACC	CTGCCCCATC	TGCCGGGATC	60
CCGCCTCGCC	CGGAAAACCG	CAGGTGACGC (GCCGACGTGG	GACGGGGTAG	ACGGCCCTAG	
					maaaamaaam	100
ATGGTCTGCG	GCAGCCCGGG .	AGGGATGCTG	CTGCTGCGGG	CCGGGCTGCT	ACCCCACCCA	120
TACCAGACGC	CGTCGGGCCC	TCCCTACGAC	GACGACGCCC	GGCCCGACGA	ACGGGACCGA	
CCTCTCTCCC	TGCTCCGGGT	CCCCCCCCT	CGGGCTGCAG	CCTGTGAGCC	CGTCCGCATC	180
CGAGAGACGG	ACGAGGCCCA	CGGGCCCCGA	GCCCGACGTC	GGACACTCGG	GCAGGCGTAG	
CCCCTGTGCA	AGTCCCTGCC	CTGGAACATG	ACTAAGATGC	CCAACCACCT	GCACCACAGC	240
GGGGACACGT	TCAGGGACGG	GACCTTGTAC	TGATTCTACG	GGTTGGTGGA	CGTGGTGTCG	
	ACGCCATCCT	GGGGN MGGN G	03 CMMCC3 3 C	CTCTCCTCCC	CACCCACTGC	300
ACTCAGGCCA	ACGCCATCCT	GGCCATCGAG	CAGTTCGAAG	GICIGCIGGG	cmcccmc100	500
TGAGTCCGGT	TGCGGTAGGA	CCGGTAGCTC	GTCAAGCTTC	CAGACGACCC	GIGGGIGACG	
オ クククククラス かく	TGCTCTTCTT	CCTCTCTCCC	ATGTACGCGC	CCATCTGCAC	CATTGACTTC	360
MOCCCCOMIC	ACGAGAAGAA	CCACACACCC	TACATCCCCC	CCTAGACCTG	GTAACTGAAG	
TCGGGGCTAG	ACGAGAAGAA	GGAGACACGG	IACATOCOCO	001110110010	4 - 1 - 1 - 1 - 1 - 1 - 1 - 1 - 1 - 1 - 	
CAGCACGAGC	CCATCAAGCC	CTGTAAGTCT	GTGTGCGAGC	GGGCCCGGCA	GGGCTGTGAG	420
GTCGTGCTCG	GGTAGTTCGG	GACATTCAGA	CACACGCTCG	CCCGGGCCGT	CCCGACACTC	
					CC1 CCTCCC1	480
CCCATACTCA	TCAAGTACCG	CCACTCGTGG	CCGGAGAACC	TGGCC'IGCGA	GGAGCTGCCA	400
GGGTATGAGT	AGTTCATGGC	GGTGAGCACC	GGCCTCTTGG	ACCGGACGCT	CCTCGACGGT	
GMGM3 GG3 G3	GGGGCGTGTG	C & MCMCMCCC	CACCCCATCC	ተሞል ርጥር ርርር A	CGGAGCTGAT	540
GIGIACGACA	CCCCGCACAC	CATCICICCC	ONCOCCHACO	* NATION COCCUM	CCCTCGACTA	
CACATGCTGT	CCCCGCACAC	GTAGAGAGGG	CTCCGGTAGC	ANIGACGCCI	GCCICGICIII	
TTTCCTATGG	ATTCTAGTAA	CGGAAACTGT	AGAGGGGCAA	GCAGTGAACG	CTGTAAATGT	600
AAAGGATACC	TAAGATCATT	GCCTTTGACA	TCTCCCCGTT	CGTCACTTGC	GACATTTACA	
						660
AAGCCTATTA	GAGCTACACA	GAAGACCTAT	TTCCGGAACA	ATTACAACTA	TGTCATTCGG	660
TTCGGATAAT	CTCGATGTGT	CTTCTGGATA	AAGGCCTTGT	TAATGTTGAT	ACAGTAAGCC	
	AAGAGATAAA	G3 GM3 3 GMCC		·	CGACCTGAAG	720
GCTAAAGTTA	AAGAGATAAA	GACTAAGTGC	CATGATGTGA	CIGCAGIAGI	CORCOLORRIC	
CGATTTCAAT	TTCTCTATTT	CTGATTCACG	GTACTACACT	GACGICATCA	CCICCACIIC	
ር ልር ልጥጥርጥል ፣	A AGTCCTCTCT	GGTAAACATT	CCACGGGAC	A CTGTCAACCT	CTATACCAGC	780
CHCHY Y CYAL	TOTOTOTOTO	ССАТТТСТА	GGTGCCCTGT	r GACAGTTGG	A GATATGGTCG	
TCTGGCTGC	TCTGCCCTCC	ACTTAATGTT	AATGAGGAA	r atatcatca	r GGGCTATGAA	840
AGACCGACG	G AGACGGGAGG	G TGAATTACAA	TTACTCCTT	A TATAGTAGT	A CCCGATACTT	

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GATGAGGAAC	GTTCCAGATT	ACTCTTGGTG	GAAGGCTCTA	TAGCTGAGAA	GTGGAAGGAT	900
CTACTCCTTG	CAAGGTCTAA	TGAGAACCAC	CTTCCGAGAT	ATCGACTCTT	CACCTTCCTA	
CC3 CMCCCM3	333330mm33					
CCMCACCCAM	AAAAAGTTAA	GCGCTGGGAT	ATGAAGCTTC	GTCATCTTGG	ACTCAGTAAA	960
GCTGAGCCAT	TTTTTCAATT	CGCGACCCTA	TACTTCGAAG	CAGTAGAACC	TGAGTCATTT	
AGTGATTCTA	GCAATAGTGA	ጥጥርርልርጥርል ር	AGTCAGAAGT	CTCCCACCAA	CDCC3 3 CCCC	1000
TCACTAAGAT	CGTTATCACT	AAGGTGAGTC	TCACTCTTCA	CIGGCAGGAA	CACCEMPOCCC	1020
			ICAGICITCA	GACCGICCII	GAGCTTGGGG	
CGGCAAGCAC	GCAACTAAAT	CCCGAAATAC	AAAAAGTAAC	ACAGTGGACT	TCCTATTAAG	1080
GCCGTTCGTG	CGTTGATTTA	GGGCTTTATG	TTTTTCATTG	TGTCACCTGA	AGGATAATTC	1000
ACTTACTTGC	ATTGCTGGAC	TAGCAAAGGA	AAATTGCACT	ATTGCACATC	ATATTCTATT	1140
TGAATGAACG	TAACGACCTG	ATCGTTTCCT	TTTAACGTGA	TAACGTGTAG	TATAAGATAA	
GTTTACTATA	AAAATCATGT	GATAACTGAT	TATTACTTCT	GTTTCTCTTT	TGGTTTCTGC	1200
CAAATGATAT	TTTTAGTACA	CTATTGACTA	ATAATGAAGA	CAAAGAGAAA	ACCAAAGACG	
##C#C#C##C	MCMO3 3 CCCC	######################################				
AAGAGAGAAG	TCTCAACCCC	TTTGTAATGG	TTTGGGGGCA	GACTCTTAAG	TATATTGTGA	1260
DAADADAAA	AGAGTTGGGG	AAACATTACC	AAACCCCCCGT	CTGAGAATTC	ATATAACACT	
GTTTTCTATT	TCACTAATCA	TCACAAAAAC	TOTAL CHARGE	C220220220	33377733303	4200
CAAAAGATAA	AGTGATTAGT	ACTCTTTTTC	ACAACAAAAC	CMATAATAAT	MARTINAACA	1320
				GITATIATIA	TITAMITIGI	
TGCTGTTACC	AGAGCCTCTT	TGCTGAGTCT	CCAGATGTTA	ATTTACTTTC	TGCACCCCAA	1380
	TCTCGGAGAA					2500
TTGGGAATGC	AATATTGGAT	GAAAAGAGAG	GTTTCTGGTA	TTCACAGAAA	GCTAGATATG	1440
AACCCTTACG	TTATAACCTA	CTTTTCTCTC	CAAAGACCAT	AAGTGTCTTT	CGATCTATAC	
CCTT 11101						
	TACTCTGCCG					1500
GGAATITIGT	ATGAGACGGC	TAGATTAATG	TCGGAATAAA	AACATACGGA	AAACCCGTAA	
CTCCTCATGC	TTAGAAAGTT	CC እ እ ከጥር ተነጥ ጥ	እጥ ል እ አርርመለ አ	እ አጥርርር እ cmm	MC33CMC333	1500
	AATCTTTCAA					1560
		COLLINCIDEN	IMITICCATI	TINCCUTCAN	ACTICAGTT	
TGTCACATAG	GCAAAGCAAT	CAAGCACCAG	GAAGTGTTTA	TGAGGAAACA	ACACCCAAGA	1620
ACAGTGTATC	CGTTTCGTTA	GTTCGTGGTC	CTTCACAAAT	ACTCCTTTGT	TGTGGGTTCT	1020
TGAATTATTT	TTGAGACTGT	CAGGAAGTAA	AATAAATAGG	AGCTTAAGAA	AGAACATTTT	1680
ACTTAATAAA	AACTCTGACA	GTCCTTCATT	TTATTTATCC	TCGAATTCTT	TCTTGTAAAA	
CCCMC3 mmc3	<u> </u>	ma				
					TAGCATTCTT	1740
CGGACTAACT	CTTCGTGTTG	ACTITIGGTCA	TUGGUGAUCC	CACAATTACC	ATCGTAAGAA	
CTTTTGGCAA	ጥACAጥጥጥርልጥ	<u> </u>	መለ ለ ጥመልጥልጥ ልጥ	CACCAMMACA	GAAATGAATT	1000
	ATGTAAACTA					1800
				CICGIANICI	CITIACTTAA	
ATAACTAGAC	ATCTGCTGTT	ATCACCATAG	TTTTGTTTAA	TTTGCTTCCT	TTTAAATAAA	1860
	TAGACGACAA					
	AAAGTCAAAA					
GGGTAACCAC	TTTCAGTTTT	TTTTTTTTT	TTT			